

## STIC Biotechnology Systems Branch

### RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/501,697  
Source: Per 110  
Date Processed by STIC: 9/6/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):  
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



PCT

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/501,697

DATE: 09/06/2005

TIME: 11:27:24

Input Set : A:\pto.kd.txt

Output Set: N:\CRF4\09062005\J501697.raw

3 <110> APPLICANT: CENTER FOR GENETIC ENGINEERING AND BIOTECHNOLOGY  
 5 <120> TITLE OF INVENTION: METHOD FOR THE GROWTHING STIMULATION AND RESISTANCE TO  
 DISEASES OF AQUATIC  
 6 ORGANISMS.  
 8 <130> FILE REFERENCE: aquatic organisms  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/501,697  
 C--> 11 <141> CURRENT FILING DATE: 2004-07-16  
 13 <150> PRIOR APPLICATION NUMBER: CU 2002-0020  
 14 <151> PRIOR FILING DATE: 2002-01-24  
 16 <160> NUMBER OF SEQ ID NOS: 1  
 18 <170> SOFTWARE: PatentIn Ver. 2.1

## ERRORED SEQUENCES

20 <210> SEQ ID NO: 1  
 21 <211> LENGTH: 6  
 22 <212> TYPE: PRT  
 23 <213> ORGANISM: artificial sequence  
 25 <220> FEATURE:  
 26 <221> NAME/KEY: PEPTIDE  
 27 <222> LOCATION: (1)..(6)  
 28 <223> OTHER INFORMATION: GHRP6  
 30 <220> FEATURE:  
 31 <223> OTHER INFORMATION: Description of the Artificial Sequence: GHRP-6  
 33 <400> SEQUENCE: 1  
 E--> 34 His D-Trp Ala Trp D-Phe Lys  
 E--> 35 1 5

Does Not Comply  
 Corrected Diskette Needed

pp 1-2

delete these- do not show them in the group of amino acids;  
 however, indicate that they are "D-form" in the  
 <2207-<2237 section. e.g. <2227 2,5  
 <2237 The amino acids at these  
 locations are D-amino acids.

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/501,697

DATE: 09/06/2005  
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Input Set : A:\pto.kd.txt  
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

## VERIFICATION SUMMARY

DATE: 09/06/2005

PATENT APPLICATION: US/10/501,697

TIME: 11:27:25

Input Set : A:\pto.kd.txt

Output Set: N:\CRF4\09062005\J501697.raw

L:10 M:270 C: Current Application Number differs, Replaced Application Number  
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:34 M:333 E: Wrong sequence grouping, Amino acids not in groups!  
L:34 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:2  
L:35 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1